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RAW SEQUENCE LISTING   DATE: 04/03/2002
PATENT APPLICATION: US/09/929,546   TIME: 16:12:06

Input Set : N:\Crf3\RULE60\09929546.raw
Output Set: N:\CRF3\04032002\I929546.raw

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 1 <110> APPLICANT: Bander, Neil H.
 2 <120> TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CANCER
 3 <130> FILE REFERENCE: Lois M. Kwasigroch: BZL 242/028
 4 <140> CURRENT APPLICATION NUMBER: 09/929,546
 5 <141> CURRENT FILING DATE: 2001-08-13
 7 <150> PRIOR APPLICATION NUMBER: 09/357,708
 8 <151> PRIOR FILING DATE: 1999-07-20
10 <150> PRIOR APPLICATION NUMBER: US 08/838,682
11 <151> PRIOR FILING DATE: 1997-04-09
12 <150> PRIOR APPLICATION NUMBER: US 60/016,976
13 <151> PRIOR FILING DATE: 1996-05-06
14 <150> PRIOR APPLICATION NUMBER: US 60/022,125
15 <151> PRIOR FILING DATE: 1996-07-18
16 <160> NUMBER OF SEQ ID NOS: 21
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 391
21 <212> TYPE: DNA
22 <213> ORGANISM: Mus sp.
23 <400> SEQUENCE: 1
24      tctcctgtca ggaactgcag gtgtcctctc tgaggtccag ctgcaacagt ctggacctga      60
25      actggtgaag cctgggactt cagtgaggat atcctgcaag acttctggat acacattcac     120
26      tgaatatacc atacactggg tgaagcagag ccatggaaag agccttgagt ggattggaaa     180
27      catcaatcct aacaatggtg gtaccaccta caatcagaag ttcgaggaca aggccacatt     240
28      gactgtagac aagtcctcca gtacagccta catggagctc cgcagcctaa catctgagga     300
29      ttctgcagtc tattattgtg cagctggttg gaactttgac tactggggcc aaggcaccac     360
30      tctcacagtc tcctcagcca aaacgacacc c                                    391
32 <210> SEQ ID NO: 2
33 <211> LENGTH: 391
34 <212> TYPE: DNA
35 <213> ORGANISM: Mus sp.
36 <400> SEQUENCE: 2
37      gggtgtcgtt ttggctgagg agactgtgag agtggtgcct tggccccagt agtcaaagtt      60
38      ccaaccagct gcacaataat agactgcaga atcctcagat gttaggctgc ggagctccat     120
39      gtaggctgta ctggaggact tgtctacagt caatgtggcc ttgtcctcga acttctgatt     180
40      gtaggtggta ccaccattgt taggattgat gtttccaatc cactcaaggc tctttccatg     240
41      gctctgcttc acccagtgta tggtatattc agtgaatgtg tatccagaag tcttgcagga     300
42      tatcctcact gaagtcccag gcttcaccag ttcaggtcca gactgttgca gctggacctc     360
43      agagaggaca cctgcagttc ctagcaggag a                                    391
45 <210> SEQ ID NO: 3
46 <211> LENGTH: 123
47 <212> TYPE: PRT
48 <213> ORGANISM: Mus sp.
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49 <400> SEQUENCE: 3
50       Ser Pro Val Arg Asn Cys Arg Cys Pro Leu Gly Pro Ala Ala Thr Val
51       1               5                   10                  15
52       Trp Thr Thr Gly Glu Ala Trp Asp Phe Ser Glu Asp Ile Leu Gln Asp
53                       20                  25                  30
54       Phe Trp Ile His Ile His Ile Tyr His Thr Leu Gly Glu Ala Glu Pro
55                       35                  40                  45
56       Trp Lys Glu Pro Val Asp Trp Lys His Gln Ser Gln Trp Trp Tyr His
57                   50                  55                  60
58       Leu Gln Ser Glu Val Arg Gly Gln Gly His Ile Asp Cys Arg Gln Val
59       65                  70                  75                  80
60       Leu Gln Tyr Ser Leu His Gly Ala Pro Gln Pro Asn Ile Gly Phe Cys
61                       85                  90                  95
62       Ser Leu Leu Leu Cys Ser Trp Leu Glu Leu Leu Leu Gly Pro Arg His
63                       100                 105                 110
64       His Ser His Ser Leu Leu Ser Gln Asn Asp Thr
65                       115                 120
67 <210> SEQ ID NO: 4
68 <211> LENGTH: 130
69 <212> TYPE: PRT
70 <213> ORGANISM: Mus sp.
71 <400> SEQUENCE: 4
72       Leu Leu Ser Gly Thr Ala Gly Val Leu Ser Glu Val Gln Leu Gln Gln
73       1               5                   10                  15
74       Ser Gly Pro Glu Leu Val Lys Pro Gly Thr Ser Val Arg Ile Ser Cys
75                       20                  25                  30
76       Lys Thr Ser Gly Tyr Thr Phe Thr Glu Tyr Thr Ile His Trp Val Lys
77                       35                  40                  45
78       Gln Ser His Gly Lys Ser Leu Glu Trp Ile Gly Asn Ile Asn Pro Asn
79                   50                  55                  60
80       Asn Gly Gly Thr Thr Tyr Asn Gln Lys Phe Glu Asp Lys Ala Thr Leu
81       65                  70                  75                  80
82       Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr Met Glu Leu Arg Ser Leu
83                       85                  90                  95
84       Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Ala Gly Trp Asn Phe
85                       100                 105                 110
86       Asp Tyr Trp Gly Gln Gly Thr Thr Leu Thr Val Ser Ser Ala Lys Thr
87                       115                 120                 125
88       Thr Pro
89                       130
91 <210> SEQ ID NO: 5
92 <211> LENGTH: 125
93 <212> TYPE: PRT
94 <213> ORGANISM: Mus sp.
95 <400> SEQUENCE: 5
96       Leu Ser Cys Gln Glu Leu Gln Val Ser Ser Leu Arg Ser Ser Cys Asn
97       1               5                   10                  15
98       Ser Leu Asp Leu Asn Trp Ser Leu Gly Leu Gln Gly Tyr Pro Ala Arg
99                       20                  25                  30
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100        Leu Leu Asp Thr His Ser Leu Asn Ile Pro Tyr Thr Gly Ser Arg Ala
101            35                  40              45
102        Met Glu Arg Ala Leu Ser Gly Leu Glu Thr Ser Ile Leu Thr Met Val
103            50                  55              60
104        Val Pro Pro Thr Ile Arg Ser Ser Arg Thr Arg Pro His Leu Thr Ser
105        65                  70              75              80
106        Pro Pro Val Gln Pro Thr Trp Ser Ser Ala Ala His Leu Arg Ile Leu
107                85                  90                  95
108        Gln Ser Ile Ile Val Gln Leu Val Gly Thr Leu Thr Thr Gly Ala Lys
109                100                 105                 110
110        Ala Pro Leu Ser Gln Pro Ser Gln Pro Lys Arg His Pro
111            115                 120                 125
113 <210> SEQ ID NO: 6
114 <211> LENGTH: 345
115 <212> TYPE: DNA
116 <213> ORGANISM: Mus sp.
117 <400> SEQUENCE: 6
118        gaggtccagc tgcaacagtc tggacctgaa ctggtgaagc ctgggacttc agtgaggata    60
119        tcctgcaaga cttctggata cacattcact gaatatacca tacactgggt gaagcagagc   120
120        catggaaaga gccttgagtg gattggaaac atcaatccta caatggtgg taccacctac     180
121        aatcagaagt tcgaggacaa ggccacattg actgtagaca agtcctccag tacagcctac   240
122        atggagctcc gcagcctaac atctgaggat tctgcagtct attattgtgc agctggttgg   300
123        aactttgact actggggcca aggcaccact ctcacagtct cctca                   345
125 <210> SEQ ID NO: 7
126 <211> LENGTH: 345
127 <212> TYPE: DNA
128 <213> ORGANISM: Mus sp.
129 <400> SEQUENCE: 7
130        tgaggagact gtgagagtgg tgccttggcc ccagtagtca aagttccaac cagctgcaca    60
131        ataatagact gcagaatcct cagatgttag gctgcggagc tccatgtagg ctgtactgga   120
132        ggacttgtct acagtcaatg tggccttgtc ctcgaacttc tgattgtagg tggtaccacc   180
133        attgttagga ttgatgtttc caatccactc aaggctcttt ccatggctct gcttcaccca   240
134        gtgtatggta tattcagtga atgtgtatcc agaagtcttg caggatatcc tcactgaagt   300
135        cccaggcttc accagttcag gtccagactg ttgcagctgg acctc                   345
137 <210> SEQ ID NO: 8
138 <211> LENGTH: 115
139 <212> TYPE: PRT
140 <213> ORGANISM: Mus sp.
141 <400> SEQUENCE: 8
142        Glu Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Thr
143        1               5                   10                  15
144        Ser Val Arg Ile Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Glu Tyr
145                    20                  25                  30
146        Thr Ile His Trp Val Lys Gln Ser His Gly Lys Ser Leu Glu Trp Ile
147                35                  40                  45
148        Gly Asn Ile Asn Pro Asn Asn Gly Gly Thr Thr Tyr Asn Gln Lys Phe
149            50                  55                  60
150        Glu Asp Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
151        65                  70                  75                  80
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152      Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
153                       85              90              95
154      Ala Ala Gly Trp Asn Phe Asp Tyr Trp Gly Gln Gly Thr Thr Leu Thr
155                  100             105             110
156      Val Ser Ser
157             115
159 <210> SEQ ID NO: 9
160 <211> LENGTH: 363
161 <212> TYPE: DNA
162 <213> ORGANISM: Mus sp.
163 <400> SEQUENCE: 9
164      ttatatggag ctgatgggaa cattgtaatg acccaatctc ccaaatccat gtccatgtca      60
165      gtaggagaga gggtcacctt gacctgcaag gccagtgaga atgtggttac ttatgtttcc     120
166      tggtatcaac agaaaccaga gcagtctcct aaactgctga tatacggggc atccaaccgg     180
167      tacactgggg tccccgatcg cttcacaggc agtggatctg caacagattt cactctgacc     240
168      atcagcagtg tgcaggctga agaccttgca gattatcact gtggacaggg ttacagctat     300
169      ccgtacacgt tcggagggggg gaccaagctg gaaataaaac gggctgatgc tgcaccaact     360
170      gta                                                                  363
172 <210> SEQ ID NO: 10
173 <211> LENGTH: 363
174 <212> TYPE: DNA
175 <213> ORGANISM: Mus sp.
176 <400> SEQUENCE: 10
177      tacagttggt gcagcatcag cccgttttat ttccagcttg gtccccccctc cgaacgtgta     60
178      cggatagctg taaccctgtc cacagtgata atctgcaagg tcttcagcct gcacactgct    120
179      gatggtcaga gtgaaatctg ttgcagatcc actgcctgtg aagcgatcgg ggaccccagt    180
180      gtaccggttg gatgccccgt atatcagcag tttaggagac tgctctggtt tctgttgata    240
181      ccaggaaaca taagtaacca cattctcact ggccttgcag gtcaaggtga ccctctctcc    300
182      tactgacatg gacatggatt tgggagattg ggtcattaca atgttcccat cagctccata    360
183      taa                                                                  363
185 <210> SEQ ID NO: 11
186 <211> LENGTH: 121
187 <212> TYPE: PRT
188 <213> ORGANISM: Mus sp.
189 <400> SEQUENCE: 11
190      Leu Tyr Gly Ala Asp Gly Asn Ile Val Met Thr Gln Ser Pro Lys Ser
191      1               5              10              15
192      Met Ser Met Ser Val Gly Glu Arg Val Thr Leu Thr Cys Lys Ala Ser
193                  20              25              30
194      Glu Asn Val Val Thr Tyr Val Ser Trp Tyr Gln Gln Lys Pro Glu Gln
195              35              40              45
196      Ser Pro Lys Leu Leu Ile Tyr Gly Ala Ser Asn Arg Tyr Thr Gly Val
197          50              55              60
198      Pro Asp Arg Phe Thr Gly Ser Gly Ser Ala Thr Asp Phe Thr Leu Thr
199      65              70              75              80
200      Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Asp Tyr His Cys Gly Gln
201                  85              90              95
202      Gly Tyr Ser Tyr Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile
203                  100             105             110
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204        Lys Arg Ala Asp Ala Ala Pro Thr Val
205            115                 120
207 <210> SEQ ID NO: 12
208 <211> LENGTH: 114
209 <212> TYPE: PRT
210 <213> ORGANISM: Mus sp.
211 <400> SEQUENCE: 12
212        Tyr Met Glu Leu Met Gly Thr Leu Pro Asn Leu Pro Asn Pro Cys Pro
213        1               5                 10                  15
214        Cys Gln Glu Arg Gly Ser Pro Pro Ala Arg Pro Val Arg Met Trp Leu
215                    20                  25                  30
216        Leu Met Phe Pro Gly Ile Asn Arg Asn Gln Ser Ser Leu Leu Asn Cys
217                35                  40                  45
218        Tyr Thr Gly His Pro Thr Gly Thr Leu Gly Ser Pro Ile Ala Ser Gln
219            50                  55                  60
220        Ala Val Asp Leu Gln Gln Ile Ser Leu Pro Ser Ala Val Cys Arg Leu
221        65                  70                  75                  80
222        Lys Thr Leu Gln Ile Ile Thr Val Asp Arg Val Thr Ala Ile Arg Thr
223                    85                  90                  95
224        Arg Ser Glu Gly Gly Pro Ser Trp Lys Asn Gly Leu Met Leu His Gln
225                    100                 105                 110
226        Leu Tyr
228 <210> SEQ ID NO: 13
229 <211> LENGTH: 116
230 <212> TYPE: PRT
231 <213> ORGANISM: Mus sp.
232 <400> SEQUENCE: 13
233        Ile Ile Trp Ser Trp Glu His Cys Asn Asp Pro Ile Ser Gln Ile His
234        1               5                 10                  15
235        Val His Val Ser Arg Arg Glu Gly His Leu Asp Leu Gln Gly Gln Glu
236                    20                  25                  30
237        Cys Gly Tyr Leu Cys Phe Leu Val Ser Thr Glu Thr Arg Ala Val Ser
238                35                  40                  45
239        Thr Ala Asp Ile Arg Gly Ile Gln Pro Val His Trp Gly Pro Arg Ser
240            50                  55                  60
241        Leu His Arg Gln Trp Ile Cys Asn Arg Phe His Ser Asp His Gln Gln
242        65                  70                  75                  80
243        Cys Ala Gly Arg Pro Cys Arg Leu Ser Leu Trp Thr Gly Leu Gln Leu
244                    85                  90                  95
245        Ser Val His Val Arg Arg Gly Asp Gln Ala Gly Asn Lys Thr Gly Cys
246                    100                 105                 110
247        Cys Thr Asn Cys
248            115
250 <210> SEQ ID NO: 14
251 <211> LENGTH: 321
252 <212> TYPE: DNA
253 <213> ORGANISM: Mus sp.
254 <400> SEQUENCE: 14
255        aacattgtaa tgacccaatc tcccaaatcc atgtccatgt cagtaggaga gagggtcacc    60
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